REMARKS

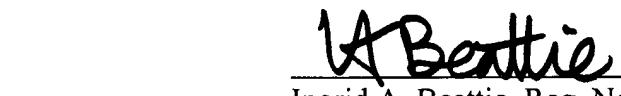
In response to the Restriction Requirement dated April 17, 2006, Applicants elect the invention of Group I, claims 1, 3, and 13-16, drawn to a method of inhibiting osteoclast-mediated bone resorption comprising inhibiting activity of a gene product encoded by an osteoclast-associated gene, and Applicants further elect the osteoclast-associated gene/marker of OC14, disclosed in Figure 1A, without traverse.

Upon entry of the present amendment, claims 1 and 25-33 are pending. Claims 2-24 have been cancelled without prejudice or disclaimer. Applicants reserve the right to prosecute the cancelled subject matter, as well as the originally presented claims, in continuing applications. Claim 1 has been amended and claims 25-33 have been added. Support for the amendments presented herein is found throughout the specification and claims as originally filed. For example, support for the amendments to claim 1 is found at least in Figure 1A, and in publicly available Genbank Accession No. AV251613, which is referred to by name in line 14 of Figure 1A: "C80638 (AV251613 RIKEN full-length enriched, 0 day neonate head Mus musculus cDNA clone 4833432F11 3', mRNA sequence)". A copy of the NCBI Entrez Database Entry for AV251613 is enclosed herewith for the Examiner's convenience. The nucleotide sequence of OC14 (AV251613) has been assigned the next available sequence identifier, *i.e.*, SEQ ID NO:50. Support for new claims 25-26 is found at least in the first paragraph on page 49 of the as-filed specification, while support for new claims 27-33 is found at least in the second paragraph on page 49 of the as-filed specification. Accordingly, no new matter has been added by the claim amendments presented herein.

CONCLUSION

On the basis of the foregoing amendment and remarks, Applicants respectfully submit, that the pending claims are in condition for allowance. If there are any questions regarding this amendment and/or these remarks, the Examiner is encouraged to contact the undersigned at the telephone number provided below.

Respectfully submitted,



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TRA 2187487v.1

Links

1: AV251613. Reports AV251613 RIKEN fu...[gi:16387654]

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LOCUS       AV251613                668 bp    mRNA    linear   EST 24-OCT-2001
DEFINITION  AV251613 RIKEN full-length enriched, 0 day neonate head Mus
            musculus cDNA clone 4833432F11 3', mRNA sequence.
ACCESSION   AV251613
VERSION     AV251613.2  GI:16387654
KEYWORDS    EST.
SOURCE      Mus musculus (house mouse)
  ORGANISM  Mus musculus
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
            Sciurognathi; Muroidea; Muridae; Murinae; Mus.
REFERENCE   1  (bases 1 to 668)
  AUTHORS   Arakawa,T., Carninci,P., Fukuda,S., Furuno,M., Hanagaki,T., Hara,A.
            , Hiramoto,K., Hori,F., Ishii,Y., Ito,M., Kawai,J., Konno,H., Kouda
            ,M., Koya,S., Matsuyama,T., Miyazaki,A., Nomura,K., Ohno,M.,
            Okazaki,Y., Okido,T., Saito,R., Sakai,C., Sakai,K., Sano,H., Sasaki
            ,D., Shibata,K., Shinagawa,A., Shiraki,T., Sogabe,Y., Suzuki,H.,
            Tagami,M., Tagawa,A., Takahashi,F., Takeda,Y., Tanaka,T., Toya,T.,
            Muramatsu,M. and Hayashizaki,Y.
  TITLE     RIKEN Mouse ESTs (Arakawa,T., et al. 2001)
  JOURNAL   Unpublished (2001)
COMMENT     On Nov 4, 1999 this sequence version replaced gi:6239072.
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            Email: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/
            Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh
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Hayashizaki,Y.
Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences Mamm. Genome. 12, 673-677 (2001)
Please visit our web site (http://genome.gsc.riken.go.jp/) for further details.
cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues.

FEATURES Location/Qualifiers
source 1..668
/organism="Mus musculus"
/mol_type="mRNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="4833432F11"
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/lab_host="DH10B"
/clone_lib="RIKEN full-length enriched, 0 day neonate head"
/note="Site_1: SalI; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was primed with a primer [5' GAGAGAGAGAAGGATCCAAGAGCTCTTTTTTTTTTTTTTTVN 3'], cDNA was prepared by using trehalose thermo-activated reverse transcriptase and subsequently enriched for full-length by cap-trapper. cDNA went through one round of normalization to Rot = 10.0 and subtraction to Rot = 100.0. Second strand cDNA was prepared with the primer adapter of sequence [5' GAGAGAGAGATTCTCGAGTTAATTAAATTAATCCCCCCCCCCCCC 3']. cDNA was cloned into the XhoI and BamHI sites. Vector: a modified pBluescript KS(+) after bulk excision from Lambda FLC I."

ORIGIN

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    481 ccgggacaac tttacttccc tttgactcag aagaaaactt cccgtggaat ttcataagca
    541 aacaaattag aaagctttac gctgctaaca gtacctcagg tgtttacttc ctcagaaaga
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    661 gattttgt
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